


IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of David V. Goeddel and Mike Rothe Serial No.: 08/446,915 Filed: 22 May 1995 For: Tumor Necrosis Factor Receptor - Associated Factors	Group Art Unit: 1812 Examiner: J. Ulm CERTIFICATE OF MAILING I hereby certify that this correspondence is being deposited with the United States Postal Service with sufficient postage as first class mail in an envelope addressed to: Assistant Commissioner of Patents, Washington, D.C. 20231 on October 3, 1997  _____ Aida A. Miclat
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PETITION TO ACCEPT PHOTOGRAPH AS DRAWING

37 CFR §1.84(b)

BOX ISSUE FEE
Assistant Commissioner of Patents
Washington, D.C. 20231

Sir:

1. Petition is hereby made to accept a photograph in this case.
2. Three (3) copies of the photograph of Figures 1, 2a, 2b, 3, 4, 5, 6a, 6b, 7, 8, 9, 15a, 15b, 16 and 17, are submitted herewith. It is submitted that photographs are the only medium by which to disclose certain aspects of the subject matter sought to be patented in this application.
3. The petition fee under 37 CFR 1.17(h) of \$130.00 is to be charged to Deposit Account No. 07-0630. Please charge any deficiency or credit any overpayment to Deposit Account No. 07-0630. A

duplicate copy of this sheet is enclosed.

Respectfully submitted,
GENENTECH, INC.

Date: October 3, 1997

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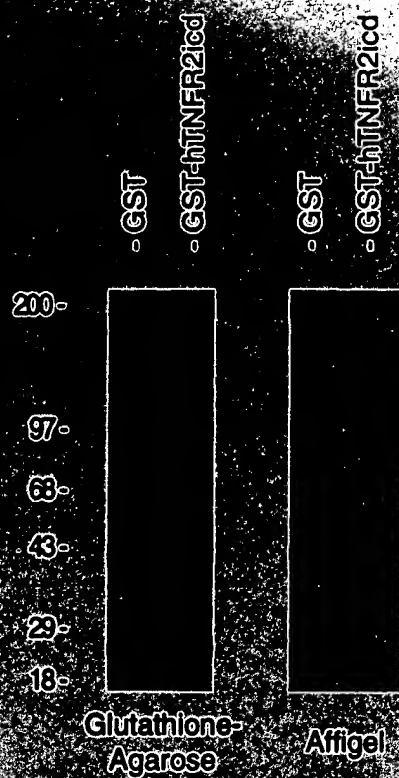


FIG. 4

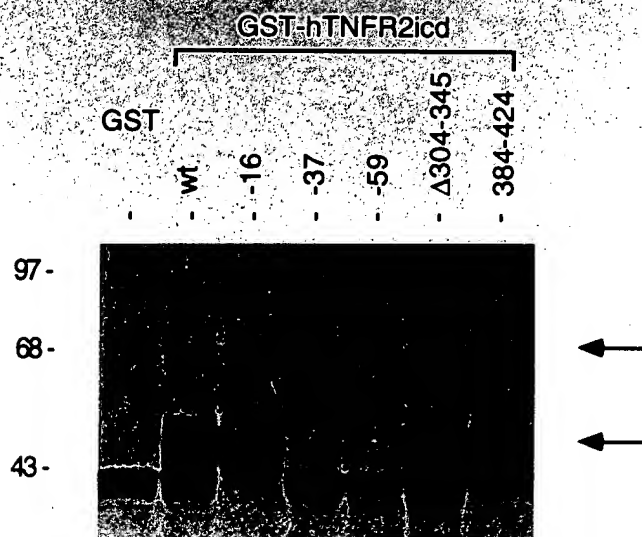


FIG. 5

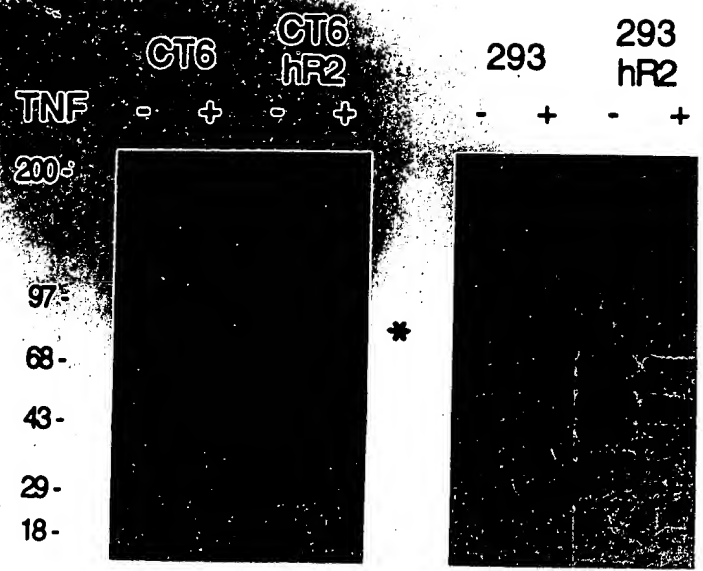


FIG. 2a

CT6
FIG. 2b

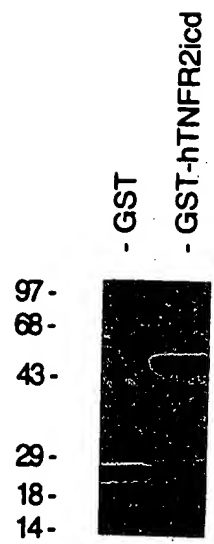


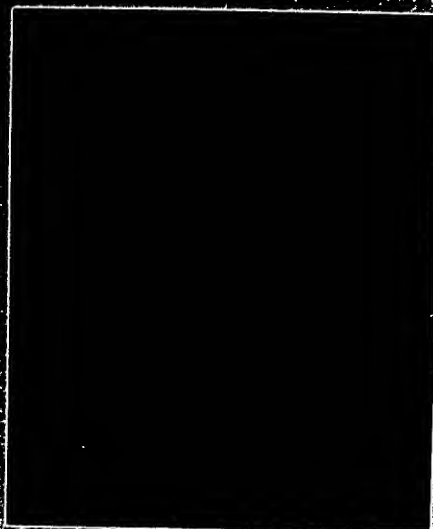
FIG. 3

5741667

Preliminary
Anti-mTNF-R2
Preliminary
Anti-mTNF-R2
Preliminary
Anti-mTNF-R2
Preliminary
Anti-mTNF-R2

NF- κ B Probe	wt	wt	mt	mt	wt	wt	wt	wt
Competitor	-	-	-	-	mt	mt	AP-1	

B



B

FIG.1

APPROVED	O.G. FIG.	
BY:	CLASS	SUBCLASS
WRAFTSMAN		

	293			293-hR2			
Competitor	-	GST	wt	-	GST	GST-hTNFR2icd	
						wt	-16 -37 -59

FIG. 6a

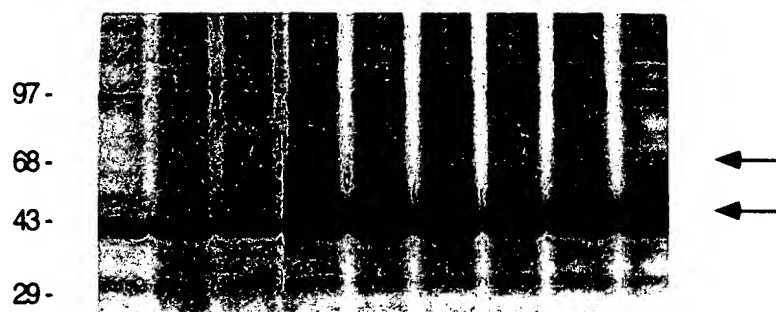
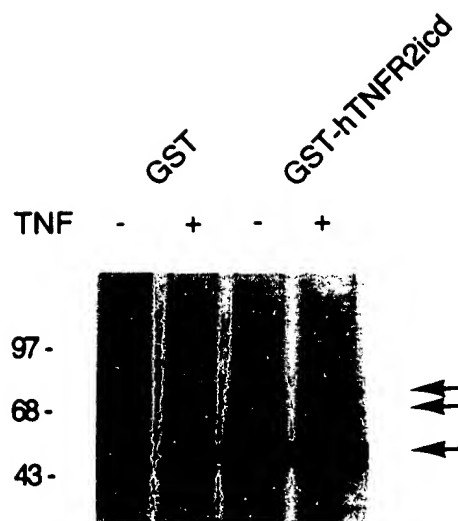


FIG. 6b



CT6

FIG. 7



ED	O.G. FIG
CLASS	SUBCLASS
DNA	MAN

		Cytoplasm					Total	
							Membrane	Extract
GST	+	-	+	-	+	-	+	-
GST-hR2icd	-	+	-	+	-	+	-	+

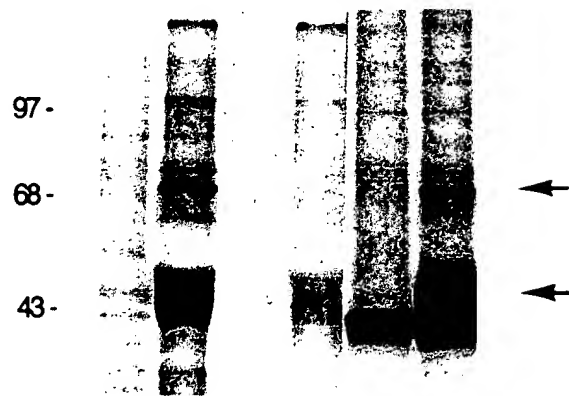


FIG. 8

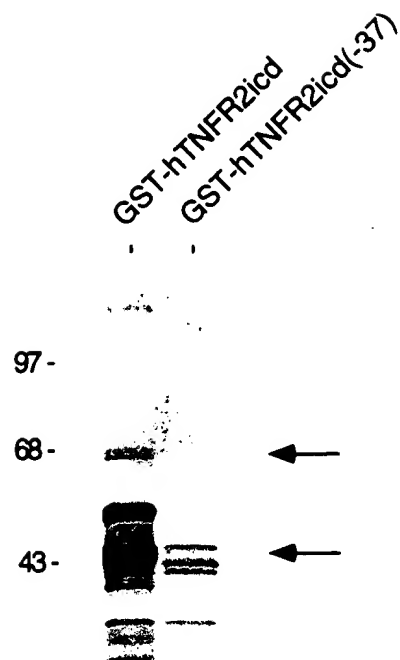


FIG. 9

APPROVED	O.G. FIG	CLASS	SUBCLASS
BY			
DRAFTSMAN			

1 CCCAGCCCGGTTCTCTGCCCCAAGGACGCTACCGCCCAATGCGAGCAGAAAGGCGCACAGATACAGAAAGT
74 GAGGCTCAGACATATTTGAGAGACCGTGTGACATAGGGTAGCCAAATGACAGTGTGAGAAAGTGACATTACTCAG
149 GCCACCCAGATATCTGGAAGGACCCAGAACCCCTGGAGATTCCCATCAGAAAGACCTCTGGCCACCTGAAACCCC
1 Meta1aSerSerSerSer1aProASD61uASng1uPheg1nPheg1uCySProPro1aProCySg1nASDPro
224 AAGATGGCTCCAGCTCAGCCCTGATGAAACGAGTTTCAATTTGGTTGCCCCCTGCTCCCTGCCAGGACCCA
25 SerG1uProArqVallleuCySCysThr1aCysleuSerG1uASnleuArqASpASD61uASPARq1leCySPro
299 TCGGAGCCAGAGTTCTCTGCTGCACAGCCTGTCTCTCTGAGAACCTGAGAGATGATGAGGATCGGATCTGTCT
50 LysCysArgAlaAspAsnleuHisProValSerProG1ySerProleuThrg1ng1uLysValHisSerAspVal
374 AAATGCAGAGCAGCAACCTCCATCTGTGAAGCCCAAGGAAGCCCTGTGACTCAGGAGAGGTTCACTCTGATGTA
75 Alag1uAlag1u1leMetCysProPhe1ag1yValG1yCysSerPhe1ysG1ySerProG1nSerMetG1ng1u
449 GCTGAAGCTGAATCATGTGCCCTTTGCAAGTGTGGCTGTTCCTTCAAGGGGAGCCCAATCCATGCAAGGAG
100 HisG1uAlaThrSerG1nSerSerHisleuTyrlleuleuAlaValleuLysG1uTrpLysSerSerProG1y
524 CATGAGGCTACCTCCAGTCTCTCCACCTGTACTGCTGCTGGCGGCTTTAAAGGAGTGGAAATCTTCACCAAGG
125 SerAsnleuG1ySer1aProMeta1aleuG1uArgASnleuSerG1uleuG1nleuG1nAla1aValG1uAla
599 TCCAACCTAGGGTCTGCACCCATGGCAC16GA6CGGAACCTGTCAAGAGCTGCAGCTTCAAGCAAGTGTGGAAGCG
150 ThrG1yASpLeuG1uValASpCysTyArqAlaProCysCysG1uSerG1ng1uG1uleuAlaleuG1nHisleu
674 ACA6GGGACCTGGA6GTAGACTGCTACCGGGACCTGTGCTGTGA6AGCCAGGAAGAACTGGCCCTGCAGCACTTG
175 ValLysG1uLysleuLeuAlag1nleuG1uG1uLysleuArqValPheAlaAsn1leValAlaValleuASnLys
749 GTGAAGGAGAGAGCTGCTGAGCTGGA6GA6AAGCTGCGTGTGTTT6CAACATTGTGTCTCAACCAAG
200 G1uValG1uAlaSerHisleuAlaleuAlaAlaSer1leHisG1nSerG1nleuASPARqG1uHisleuLeuSer
824 GAAGTGGAGGCTTCCACCTGGCAC16GCCGCTCCATCCACAGAGCCAGTTGGACCGAGAGCACTCTGAGC
225 LeuG1uG1nArqValValG1uleuG1ng1nThrleuAlag1nLysASpG1nValleuG1yLysleuG1uHisSer
899 TTGGAGCAGAGG6GTGGATTAACAGCAAAACCTGGCTCAAAAAGACCAAGGCTCTGGGCAAGCTTGAAGCACAGT

FIG. 10a

 \wedge

FIG. 10b

APPROVED	O G FIG	DRAFTSMAN
B*	CLASS	
	SUBCLASS	

* * *

1 GC GCGAAGACCGTTGGGGCTTGTGGTGTGGGGGTTGTAAC TCACATGGCTGCAGCCAGTGTGACTTCCCC
10 GlysSerLeuGluLeuLeuGlnProGlyPheSerLysThrLeuLeuGlyThrArgLeuGluAlaLysTyrLeuCys
75 GGCTCCCTAGAACTGCTACAGCTGGCTTCTCCAAGACCTCTCTGGGGACCAGGTTAGAAAGCCAAGTACCTCTGT
35 SerAlaCysLysAsnIleLeuArgArgProPheGlnAlaGlnCysGlyHisArgTyrCysSerPheCysLeuThr
150 TCAGCCTGC AAAACA TCCTGCGGAGGCCCTTCCAGGCCAGTGTGGGCACCGCTACTGCTCTTGCTGACC
60 SerIleLeuSerSerGlyProGlnAsnCysAlaAlaCysValTyrGluGlyLeuTyrGluGlyIleSerIle
225 AGCATCCTCAGCTCTGGGCCCCAGAACTGTGCTGTGTCTATGAAGGCCCTGTATGAAGAAAGCAITTTCTATT
85 LeuGluSerSerSerAlaPheProAspAsnAlaAlaArgArgGluValGluSerLeuProAlaValCysProAsn
300 TTAGAAGAGTAGTTCGGCTTTCAGATAACGCTGCCCGCAGAGAGGTGGAGAAGCTGCCAGCTGTCTGTCCCAAT
110 AspGlyCysThrTrpLysGlyThrLeuLysGluTyrGluSerCysHisGluGlyLeuCysProPheLeuLeuThr
375 GATGGATGCACCTGGAAAGGGGACCTTGAAAGAAATACGAGAAGCTGCCACGAAAGGACTTGGCCATTCTGCTGACG
135 GluCysProAlaCysLysGlyLeuValArgLeuSerGluLysGluHisThrGluGluGluCysProLysArg
450 GAGTGTCTGCATGTAAAGGCTGTGTCGCTCAGCGAGAGAGGACACACACTGAGCAGGAATGCCCAAAGG
160 SerLeuSerCysGlnHisCysArgAlaProCysSerHisValAspLeuGluValHisTyrGluValCysProLys
525 AGCTGAGCTGCCAAGCACTGCAGAGCACCTGTAGCCACGTGGACCTGGAAGTACACTATGAGGCTTGCCCCAAG
185 PheProLeuThrCysAspGlyCysGlyLysLysLysIleProArgGluThrPheGlnAspHisValArgAlaCys
600 TTTCCCTAACCTGTGATGGCTGTGGCAAGAAGAGATCCCTCGGGAGACGTTTCAAGACCATGTAGAGCAATGC
210 SerLysCysArgValLeuCysArgPheHisThrValGlyCysSerGluMetValGluThrGluAsnLeuGlnAsp
675 AGCAATGCGGGGTTCTCTGCAAGATTCACACACCGTTGGCTGTTCAGAGATGGTGGAGACTGAGAACCTGCAGGAT
235 HisGluLeuGlnArgLeuArgGluHisLeuAlaLeuLeuLeuSerSerPheLeuGluAlaGlnAlaSerProGly
750 CATGAGCTGCAGCGGCTACGGGAAACACCTAGCCCTACTGCTGAGCTCATTTCTGGAGGGCCCAAGCCTCTCAAGGA
260 ThrLeuAsnGlnValGlyProGluLeuLeuGlnArgCysGlnIleLeuGluGlnLysIleAlaThrPheGluAsn
825 ACCTTGAACCAAGGTGGGGCCAGAGCTACTCCAGCGGTGCAAGATTTTGGAGCAGAAAGATAGCAACCTTTGAGAAC

FIG. 11a

285 ILevalCysValLeuAsnArggIuValGIuArgValAlaValIThrAlagIUA1aCysSerArggIhHisArgLeu
900 ATTGCTGCGCTTGAACCGTGAAGTAGAGAGGGTAGCAGTGAAGTGAAGGCTTGTAGCCGGCAGCAGGCTA
310 AspGIhAspLysI1egIUA1aLeuSerAsnLysValGInGInLeuGIuArgSerI1egI1eulLysAspLeuAla
975 GACCAGGACAAGATTGAGGCCCTGAGTAACCAAGGTGCAACAGCTGGAGAAGGATCGGCCCTCAAAGACCTGGCC
335 MetAlaAspLeuGIuGIuLysValSerGIuLeuGIuValSerIThrAspGIuValPheI1eIrpLysI1eSer
1050 ATGGCTGACCTGGAGCAGAAAGGCTCCGAGTTGGAAAGTATCCACCTATGATGGGGCTTTCATCTGGAAAGATCTCT
360 AspPheThrArgLysArgGInGIuAlaValAlagIArgIThrProAlaI1ePheSerProAlaPheIYrThrSer
1125 GACTTCACCAGAAAGCCGTCAGGAAGCCGTAAGTGGCCGGACACCAGCTATCTTCCCCAGGCTTCTACACAAAGC
385 ArgIYrGLYIYrLysMetCysLeuArgValIYrLeuAsnGIuAspGIuThrGIuArgGIuIYrHisLeuSerLeu
1200 AGATATGGCTACAAGAGTGTGTACGAGTCTACTTGAATGGCGACGGCACTGGGCGGGGAACCTACTGTCTCTC
410 PhePheValValMetLysGIuProAsnAspAlaLeuLeuGIuNTIrpProPheAsnGIuLysValIThrLeuMetLeu
1275 TTCCTTGGTGGTGAAGAAAGGCCCAATGATGCTCTGTGGAGTGGCTTTTAATCAGAAAGTAACATTGATGTTG
435 LeuAspHisAsnAsnArgGIuHisValI1eAspAlaPheArgProAspValIThrSerSerPheGInArgPro
1350 CTGGACCATAAACAACCGGAGCATGTGATCGACGCATTCAAGGCCGATGTAACTCGTCCCTCTCAAGAGGCCCT
1425 ValSerAspMetAsnI1eAlaSerGIuYrCysProLeuPheCysProValSerLysMetGIuAlaLysAsnSerIYr
1425 GTCAGTGACATGAACATCGCCAGTGGCTGGCCCTCTTCTGCCCTGTGTCCAAGATGGAGGCCAAGAAATCTCTAT

485 ValArgAspAspAlaI1ePheI1eLysAlaI1eValAspLeuThrGIuLeu
1500 GTCCGGGATGATGCGATCTTCATCAAGCTATTGTGACCTAACAGGACTCTAGCCACCCCTGCTAAGAATAGCA
1575 GCTCAGTGAAGGAGCTGTACATTAGGCCAGCCAGCCCTGCCACACACGGGTGGCCAGGCTTGGTGTAAATGCTG
1650 GGGAGGGCTCAGCTAGAGCCAAATCACCATCACACAGAAAGGAGGAGAAAGGCTCCAGTTGGCTTCAAGTGG
1725 CAACTGAGTTGGACGGTCCACTGAGCTCAAGGGCTGGTGAAGCCGCTGGGGAAGCTTCTCAGCTTCCAATAG
1800 GAAAGCTCTGCTGTCTCTGTCTGTCTGGGGAAAGGAGAGACTGTAGGTGGGTGCTCAGAAAGGGCTCTCCAGA
1875 GAGAGTCTCAAGAGCTGACAGCAGGAGCAAAAGTGAAGTGGCTTCCCAACCCCATCTTGGAAAGAGAGGTAGCGGC
1950 TACACAGGAGAGGAGCATGGCTTGCAGGGTGTAGCCCAAGAGAGAGAGCTCTGTGAACATAGGCCCTCACTGGAG
2025 AAGGGCTTGGCTGGGCTGCACAGCCTTGGCAGGTGGCTGTATGGGGGAGAGTGAATTAATGTGTAGATGTAC
2100 ACGACAAAAAATAAAAAA

FIG. 11b

FIG. 12a

TRAF2	(mouse)	31	KYLCSAC	KNILRRPFQA	QCGHRYC	SFC	LTSI	LSS	GPQNCA	CAYE
COP1	(A. thaliana)	49	DLLCPIC	MOIIDAFLT	ACGHSFC	YMC	IIITH	LRN	KSDC	PCCSQH
EFP	(human)	10	ELSCSIC	LEPFKEPVTP	PCGHNFC	GSC	LNETWA	VQG	SPYLC	PQCR
RAD-18	(S. cerevisiae)	25	LLRCHIC	KDFLKVPLT	PCGHTFC	SLC	IRTH	LNN	QPNCP	LCLFE
UVS-2	(N. crassa)	31	AFRC	HVC	KDFYDSPMLT	SCNHTFC	SLC	IRRC	LSV	DSK
RAG-1	(human)	290	SISCQIC	EHILADPVET	MC	KHVC	RVCI	LRG	LKV	MGSYC
SS-A/Ro	(human)	13	EVTCPIC	LDPFVEPVS	ECGHSFC	QEC	ISQV	GKG	GGSV	CAVCRQR
RING1	(human)	16	ELMCPIC	LDMLKNTMTTKECL	HRFC	SDC	IVTA	LRS	GNKEC	PTCRKX
RPT-1	(mouse)	12	EVTCPIC	LELKEPVSA	DCNHSFC	RAC	ITLNYESN	RNTDGKGN	CPVC	RVVP
RFP	(human)	13	ETT	CPVCLQYFAEP	PML	DCGHNIC	CA	LARCW	GTA	ETNVS
c-cbl	(human)	378	FQLCKI	CAENDKDVKIE	PCGHL	MC	TSC	LTS	WQESGQ	GSSGCP
consensus										

-----X11-12-----C-H-C-----X10-16-----C-C-----

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

FIG. 12b

TRAF2	(mouse)	157	CPKRSLS C QH C RAPCSHVDLEV H YE V C
		182	PKFPLT C DG C GKKKIPRET F QD H VR A C
DG17	(<i>D. discoideum</i>)	171	GGFKLVT C DF C KRDDIKKKELET H YK T C
TFIIIA	(<i>X. laevis</i>)	189	QD LAV C DV C NRKFRHKDYLRD H QK T H
XLCOF14	(<i>X. laevis</i>)	1	TGKYPFI C SE C GKSFMDKRYLKI H SN V H
XFIN	(<i>X. laevis</i>)	1225	TGEKPYT C TV C GKKFIDRSSVVR H SR T H
ZFY1/2	(mouse)	521	RKKFPHI C GE C GKGFRHPSALKK H IR V H
MFG2	(mouse)	293	SEEKPF E CE E GKKFRTARHLVR H QR I H
RAD18	(<i>S. cerevisiae</i>)	183	PNEQMAQ C PI C QQFYPLKALEKT H LD E C
UVS-2	(<i>N. crassa</i>)	182	PDDGLVA C PI C LTRM KEQQVDR H LDTS C

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

TRAF2 1 MAAASVTSPGSLLELLQPGFSKTLTGTRLEAKYLC SACKNILRRPFOAQCG

TRAF2 51 HRYCSFCLTSILSSGPONCAACVYEGLYEEGISILESSSAFPD NAARREV

TRAF2 101 ESLPAVCPNDGCTWKGT LKEYESCHEGLCPFLLTECPACKGLVRLSEKEH

TRAF1 1 MASSAPDENE FQFGCPPA

TRAF2 151 HTEQECPKRSLSCOHCRAPCSHVDLEVHYEVC PKFPLTCDGC GKKKIPRE

TRAF1 20 PCQDPSEPRVLCCTACLSENLRDDEDRICPKCRADNLHPVSPG SPTLQE

TRAF2 201 TFQDHVRACSKCRVLCRFHTVGCSSEMVETENLQDH E LQRLREHLALLSS

TRAF1 69 KVHSDV . . . AEAEIMC PFAGVGCSFKGSPQSMQEHEATSQSSHL Y LLLAV

TRAF2 251 FLEAQA SPGTLNQVGPELLQR

TRAF1 116 LKEWKS SPGSLGSA PMALERNLSELQLQA AVEATGDLEVDCYRAPCCES

TRAF2 272 CQILEQKIATFENIVCVLNREVERVAVTAEACSRQH

TRAF1 166 QEELALQHLVKEKLLAQLEEKLRVFANIVA VLNKEVEASHLALAASIHQS

TRAF2 308 RLDQDKIEALSNKVQQLERSIGLKDLAMADLEQKVSEL EVSTYDGVFIWK

TRAF1 216 QLDREHLLSLEQRVVELQQT LAQKDQVLGKLEHSLRLMEEASF DGTFLWK

TRAF2 358 ISDFTRKRQEA VAGRTPAIFSPA FYTSRYGYKMCLRVYLN GDGTGRGTHL

TRAF1 266 ITNVTKRCHESVCGRTVSLFSPA FYTAKYGYKLCRL YLNGDGS GK KTHL

TRAF2 408 SLFFVVMKGPNDALLQWPFNQKVTLMLLDHNNREHV IDAFRPDVTSSSFQ

TRAF1 316 SLFIVIMRGEYDALLPWPFRNKVTFMLLDQNNREH IDAFRPDLS SASFQ

TRAF2 458 RPIVSDMNIASGCPLFCPVSKME AKNSYVRDDAIFIKAI VDLTGL

TRAF1 366 RPQSETNIVASGCPLFFPLSKLQSPKHAYVKDDT MFLKCI VDTSA

FIG. 13

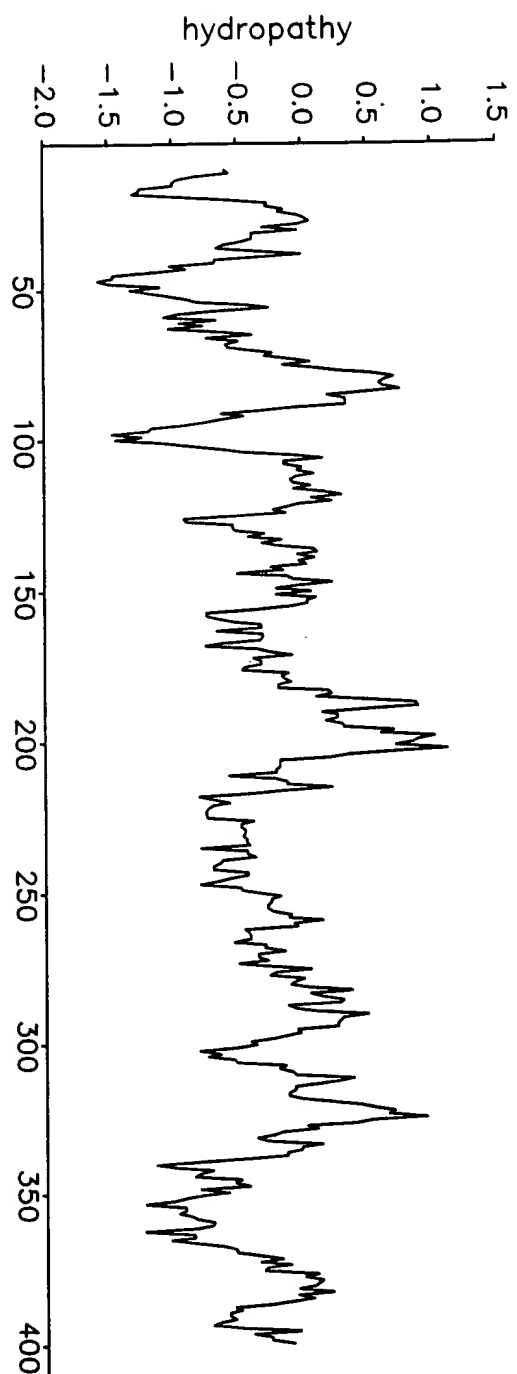


FIG. 14a

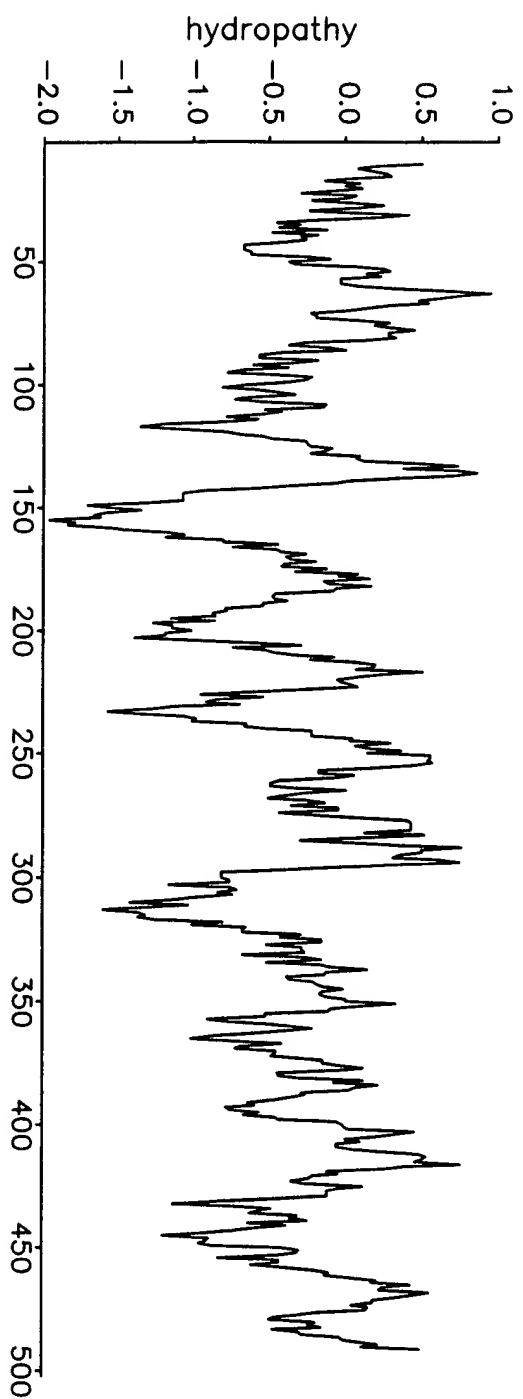


FIG. 14b

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

FIG. 18a

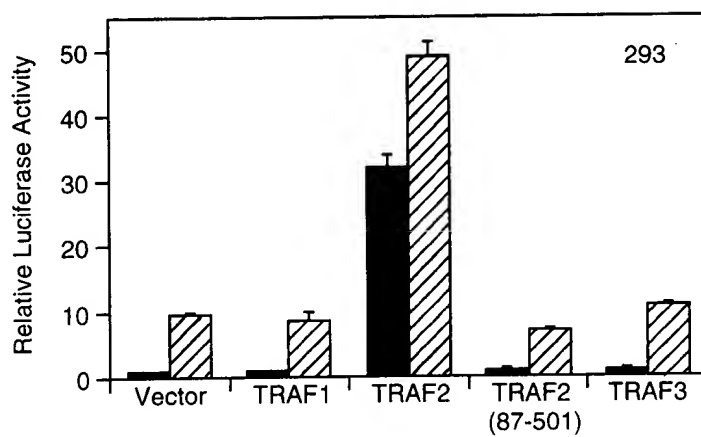
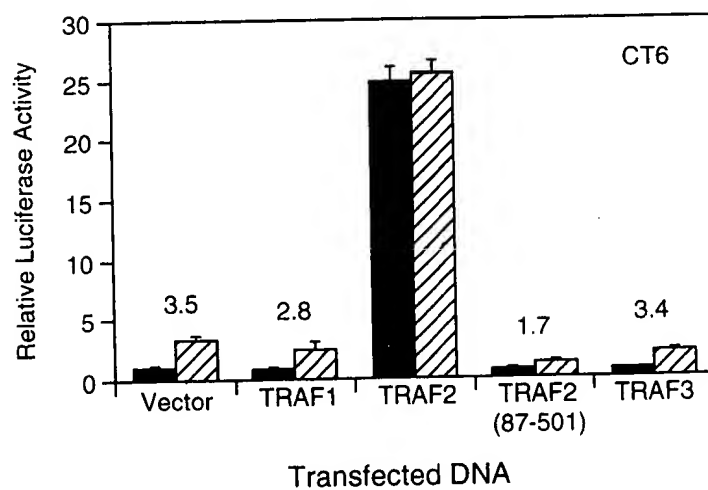


FIG. 18b



APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

FIG. 19a

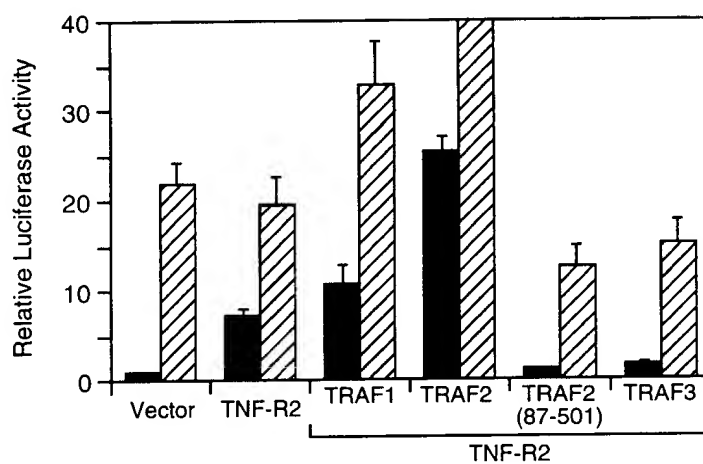
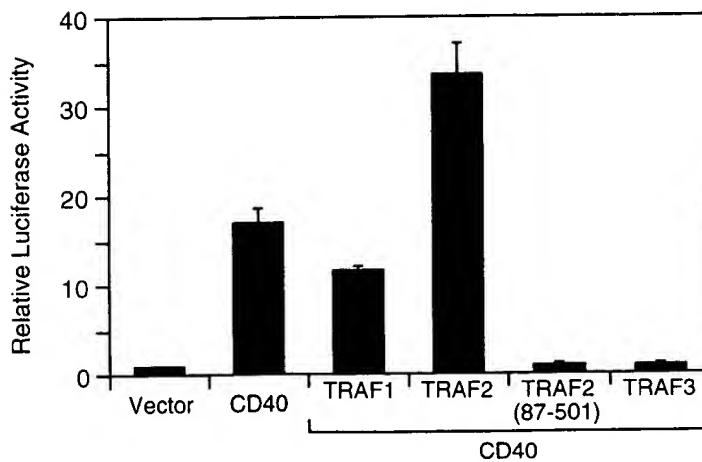


FIG. 19b



Transfected DNA

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

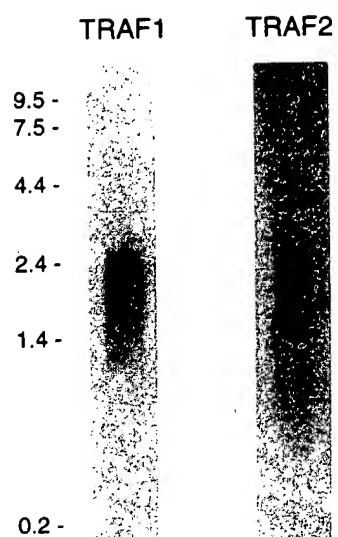


FIG. 15a

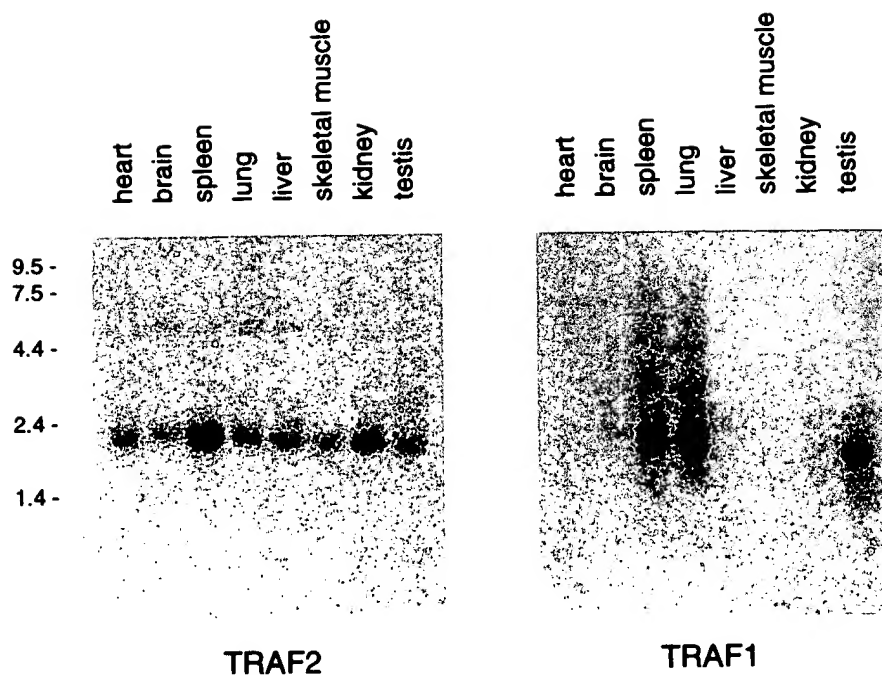


FIG. 15b

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

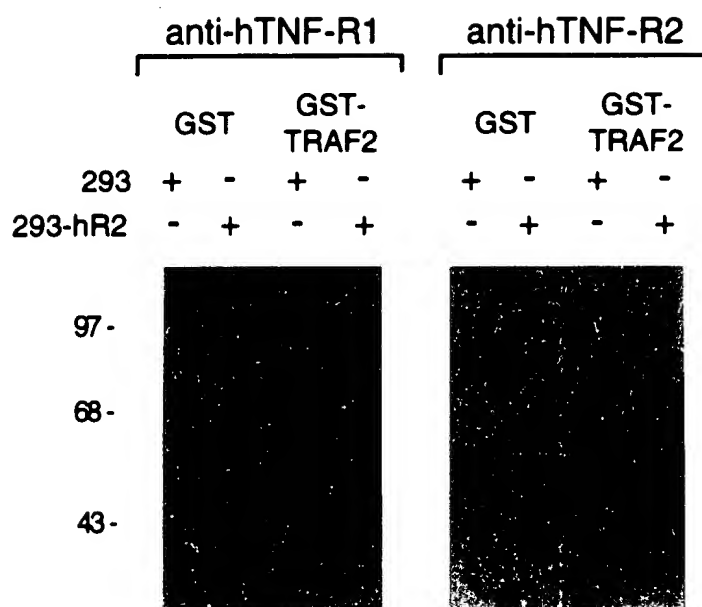
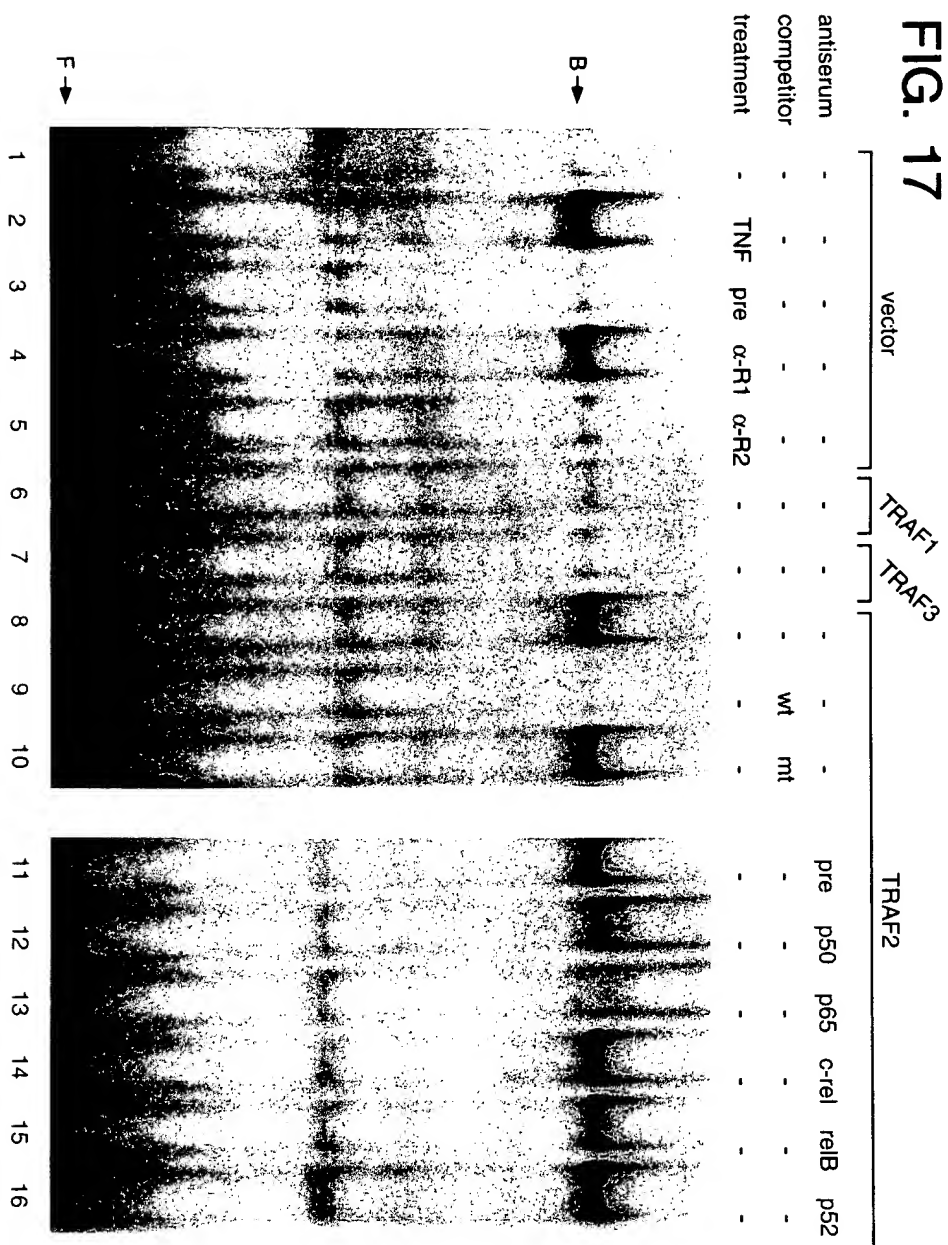


FIG. 16

FIG. 17



O.G. FIG.		APPROVED	DRAFT
CLASS		SUBCLASS	